

SEQUENCE LISTING



#10/10

<110> ASHKENAZI, AVI J.  
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 WOOD, WILLIAM I.  
 GENENTECH INC.

<120> DcR3 Polypeptide, A TNFR Homolog

<130> 11669.31US03

<140> 09/157,289

<141> 1998-09-18

<150> 60/059,288

<151> 1997-09-18

<150> 60/094,640

<151> 1998-07-30

<160> 16

<170> PatentIn Ver. 2.0

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Thr	Pro	Thr	Tyr	Pro	Trp	Arg	Asp	Ala	Glu	Thr	Gly	Glu	Arg	Leu	Val
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Cys	Ala	Gln	Cys	Pro	Pro	Gly	Thr	Phe	Val	Gln	Arg	Pro	Cys	Arg	Arg
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Asp	Ser	Pro	Thr	Thr	Cys	Gly	Pro	Cys	Pro	Pro	Arg	His	Tyr	Thr	Gln	65	70	75	80
Phe	Trp	Asn	Tyr	Leu	Glu	Arg	Cys	Arg	Tyr	Cys	Asn	Val	Leu	Cys	Gly	85	90	95	
Glu	Arg	Glu	Glu	Glu	Ala	Arg	Ala	Cys	His	Ala	Thr	His	Asn	Arg	Ala	100	105	110	
Cys	Arg	Cys	Arg	Thr	Gly	Phe	Phe	Ala	His	Ala	Gly	Phe	Cys	Leu	Glu	115	120	125	
His	Ala	Ser	Cys	Pro	Pro	Gly	Ala	Gly	Val	Ile	Ala	Pro	Gly	Thr	Pro	130	135	140	
Ser	Gln	Asn	Thr	Gln	Cys	Gln	Pro	Cys	Pro	Pro	Gly	Thr	Phe	Ser	Ala	145	150	155	160
Ser	Ser	Ser	Ser	Ser	Glu	Gln	Cys	Gln	Pro	His	Arg	Asn	Cys	Thr	Ala	165	170	175	
Leu	Gly	Leu	Ala	Leu	Asn	Val	Pro	Gly	Ser	Ser	Ser	His	Asp	Thr	Leu	180	185	190	
Cys	Thr	Ser	Cys	Thr	Gly	Phe	Pro	Leu	Ser	Thr	Arg	Val	Pro	Gly	Ala	195	200	205	
Glu	Glu	Cys	Glu	Arg	Ala	Val	Ile	Asp	Phe	Val	Ala	Phe	Gln	Asp	Ile	210	215	220	
Ser	Ile	Lys	Arg	Leu	Gln	Arg	Leu	Leu	Gln	Ala	Leu	Glu	Ala	Pro	Glu	225	230	235	240
Gly	Trp	Gly	Pro	Thr	Pro	Arg	Ala	Gly	Arg	Ala	Ala	Leu	Gln	Leu	Lys	245	250	255	
Leu	Arg	Arg	Arg	Leu	Thr	Glu	Leu	Leu	Gly	Ala	Gln	Asp	Gly	Ala	Leu	260	265	270	
Leu	Val	Arg	Leu	Leu	Gln	Ala	Leu	Arg	Val	Ala	Arg	Met	Pro	Gly	Leu	275	280	285	
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aggcctgtcg ctgctgtgcc tgggtgttggc gctgcctgcc ctgctgccgg tgcgggctgt 180  
acgcggagtg gcagaaacac ccacctaccc ctggcgggac gcagagacag gggagcggct 240  
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cacgacgtgt ggcccgtgtc caccgcgcc aacacgcagc ttctggaact acctggagcg 360  
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caccacaac cgtgcctgcc gctgcgcgac cggtctcttc gcgcacgctg gtttctgctt 480  
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cacgcagtgc cagccgtgcc cccaggcac cttctcagcc agcagctcca gctcagagca 600  
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gggcttgcca cgccaccac aaccgtgcct gccgctgccg caccggcttc ttcgcgcacg 180
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ccccagcca gaacacgcag tgcctagccg tgcctccag gcaccttctc agccagcagc 300
tccagctcag agcagtcca gcccaccgc aactgcacgg ccctgggcct ggccctcaat 360
gtgccaggct cttcctcca tgacaccctg tgcaccagct gcactggctt cccctcagc 420
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gngcttgcca cgccaccac aaccgcgcct gcnctgcag caccgnttc ttgcgcacg 180
ctgntttctg cttggagcac gcatcgtgtc cacctggtgn cggcgtgatt gcncgggca 240
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cttcgcgcac gctggtttct gcttgagca cgcctcgtgt ccacctggtg ccggcgtgat 180
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acccccagcc agaacacgca gnccagccgt gccccagcag caccttctca gccagcagct 180
ccagctcaga gcagtgccag cccacccgca actgcacggc cctgggcctg gccctcaatg 240
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cgcaactgca acgccctggn ctggccctca atgtgccagg ctcttctctc catgacaccc 180
tgtgcaccag ctgcactggc ttccccctca gcaccagggt accaggagct gaggagtgtg 240
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22

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17

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